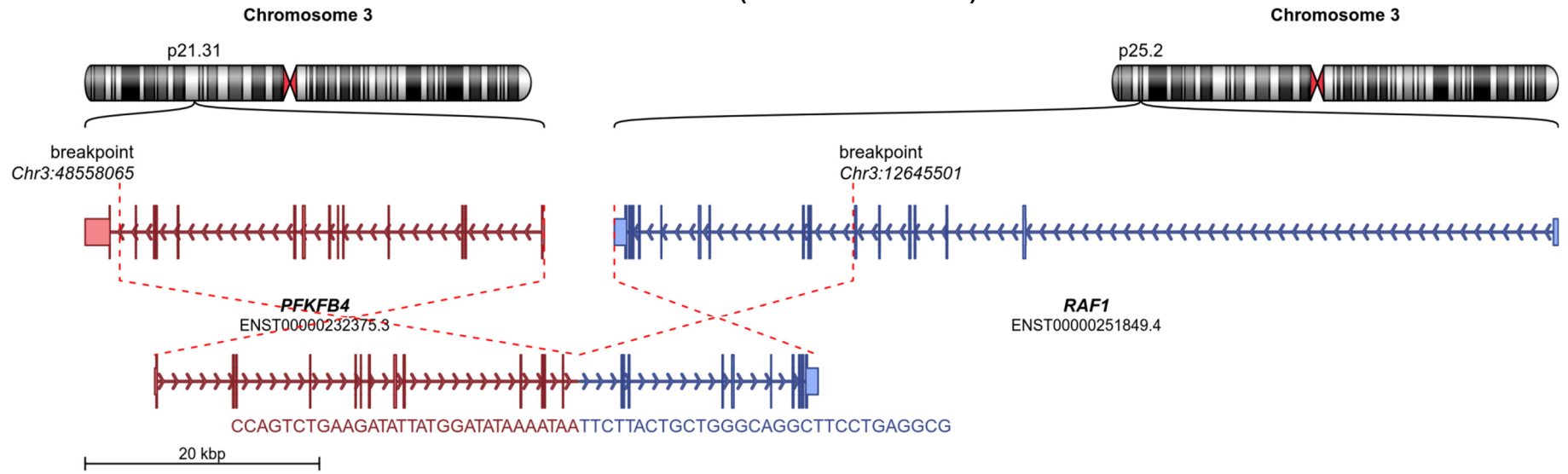
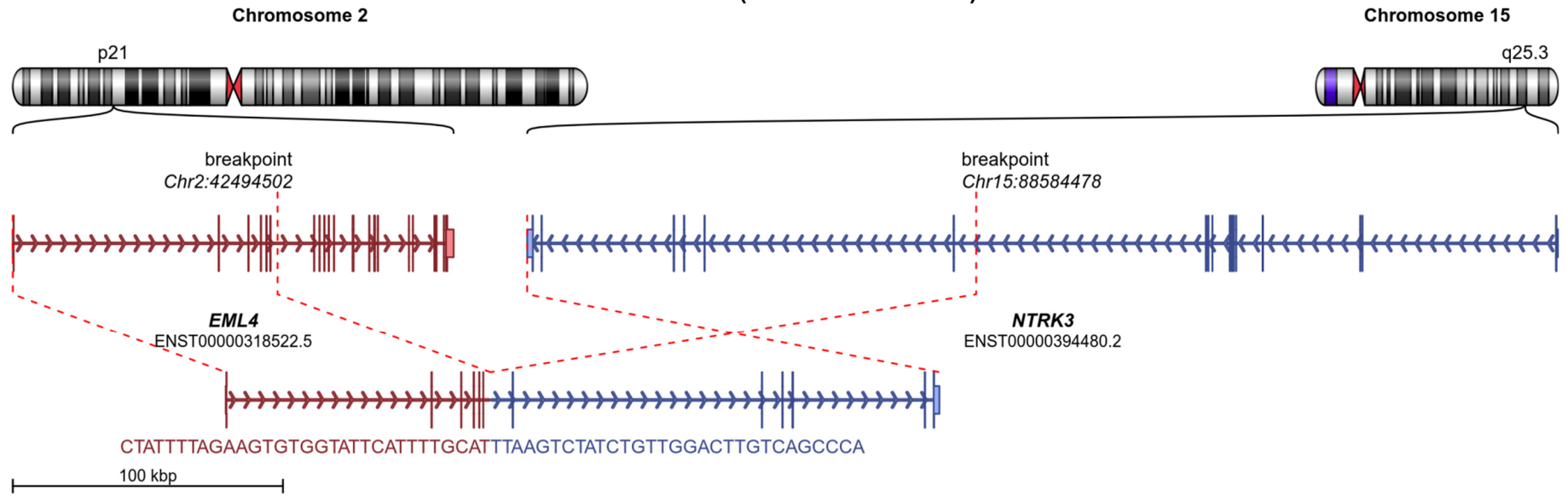


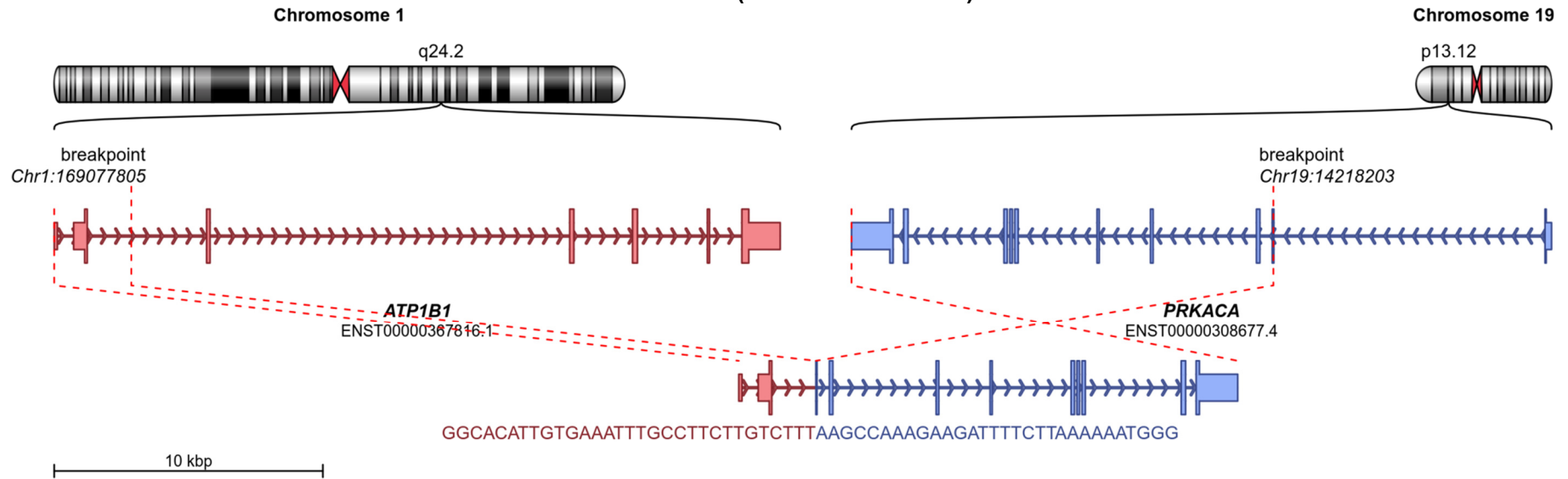
COMP-002 (EGAF00001709816)



COMP-0023 (EGAF00001709871)



COMP-0028 (EGAF00001709880)



COMP-0055 (EGAF00001720843)

Chromosome 20



p12.1

breakpoint
Chr20:17611430



RRBP1
ENST00000377807:2

Chromosome 3



p25.2

breakpoint
Chr3:12657584

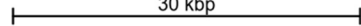


RAF1
ENST00000251849.4

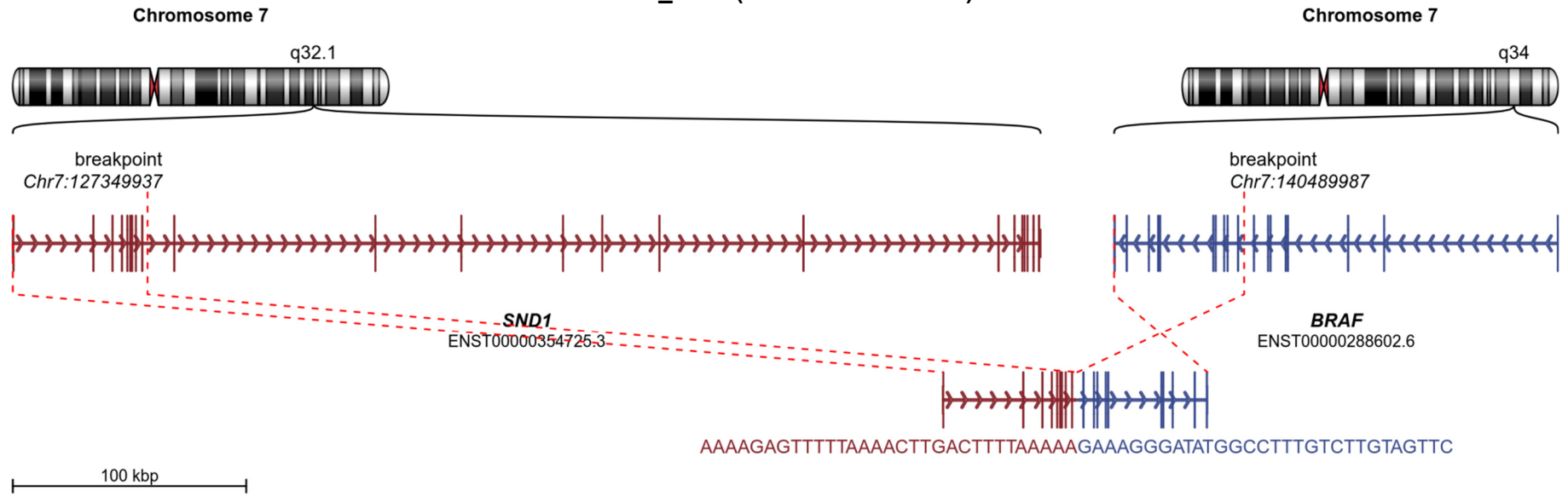


GGTGTGGTGGGCCTGTCTGTCTGAGTGAGGTAGCTATTATTATTATTATTATGATTT

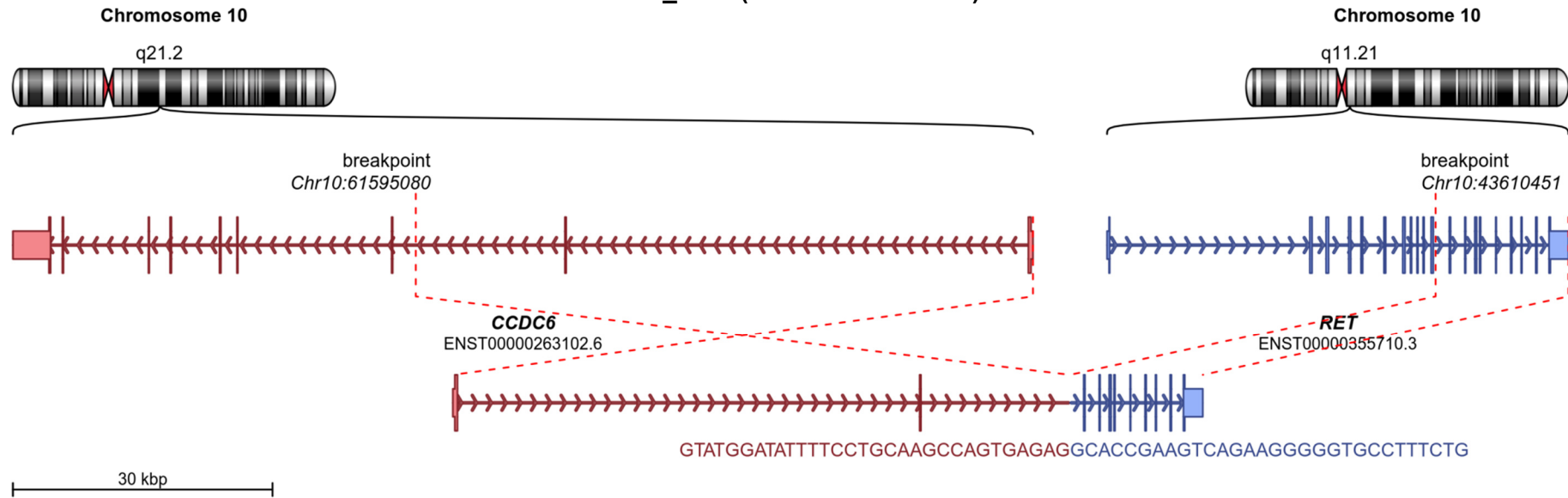
30 kbp



ICGC_0338 (EGAF00001144272)



ICGC_0391 (EGAF00001144341)



Patient 14 (EGAF00001984740)

Chromosome 1

Chromosome 8

q24.2

p12

breakpoint
Chr1:169088266

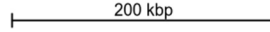
breakpoint
Chr8:32447498

ATP1B1
ENST00000367816.1

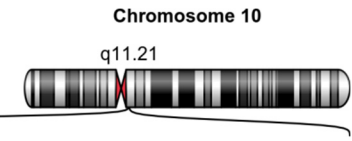
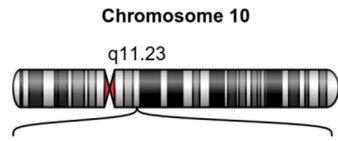
NRG1
ENST00000520407.1

GAAATGATGAACCTTGCTACATTATAAATTGTGTGACATAGCAAATAGGAAAAGCAGA

200 kbp

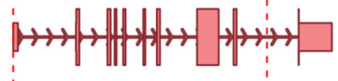


Patient 17 (EGAF00001984732)



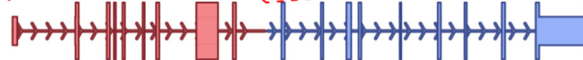
breakpoint
Chr10:51587787

breakpoint
Chr10:43611343



NCOA4
ENST00000443446.1

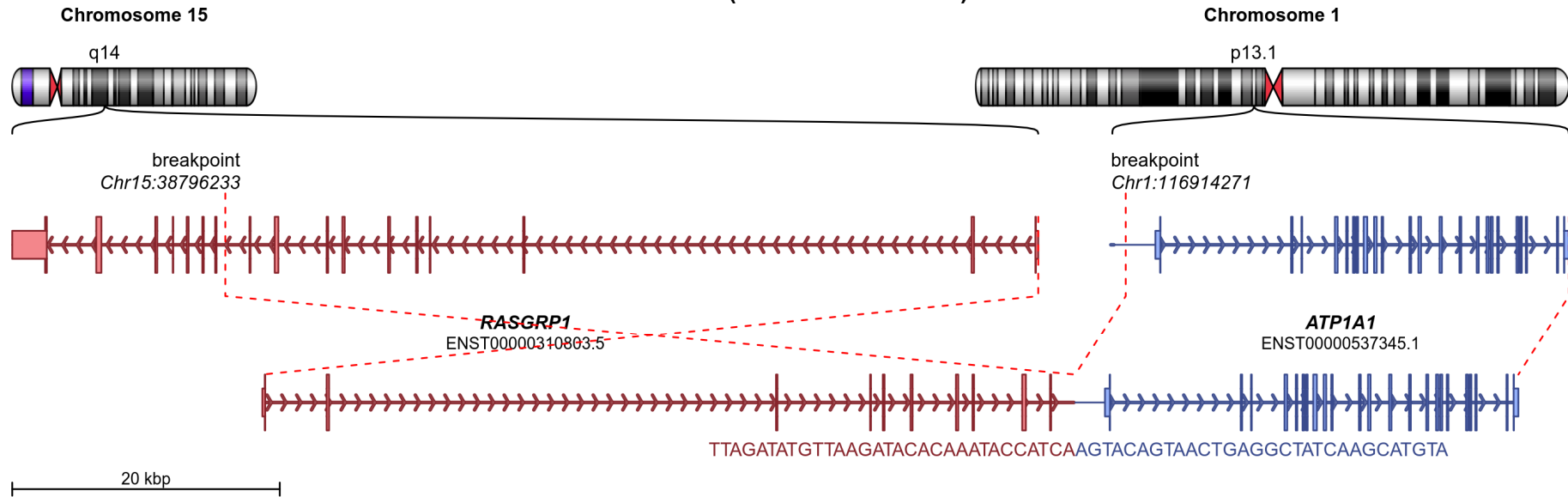
RET
ENST00000355710:3



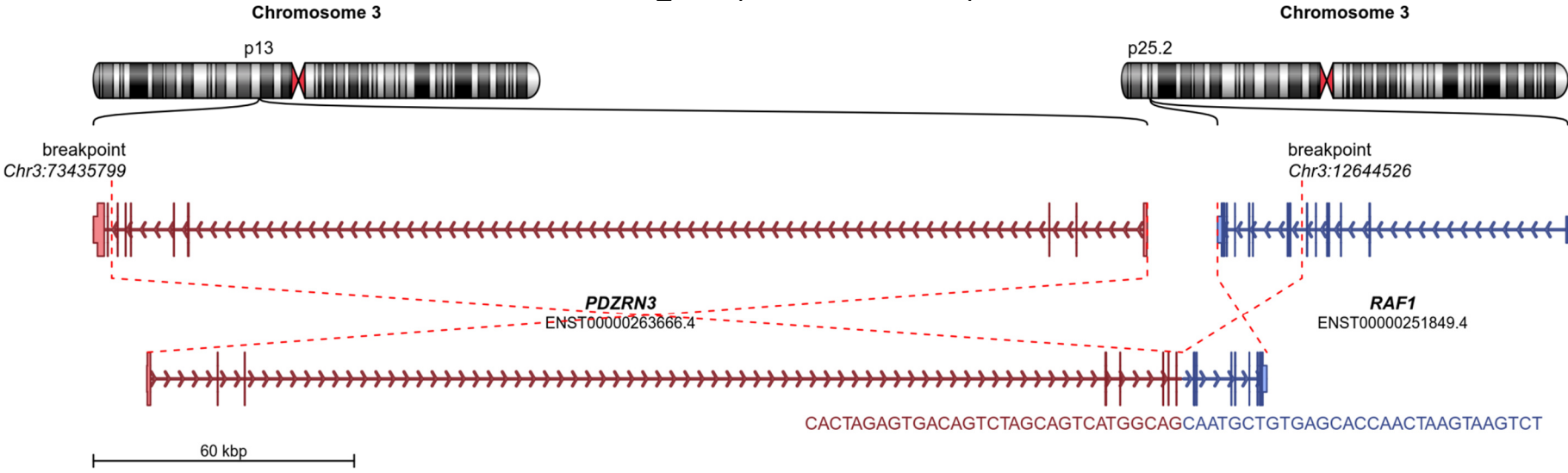
CAGTAGTTCTCAACAGAGGGCTTAAAATATGCTAAATCTACTCTGCCTTGTGCTCTAGAAATGGAAAGGGGCCAGACCAGGACAGCCCAGGAGCAGG

10 kbp

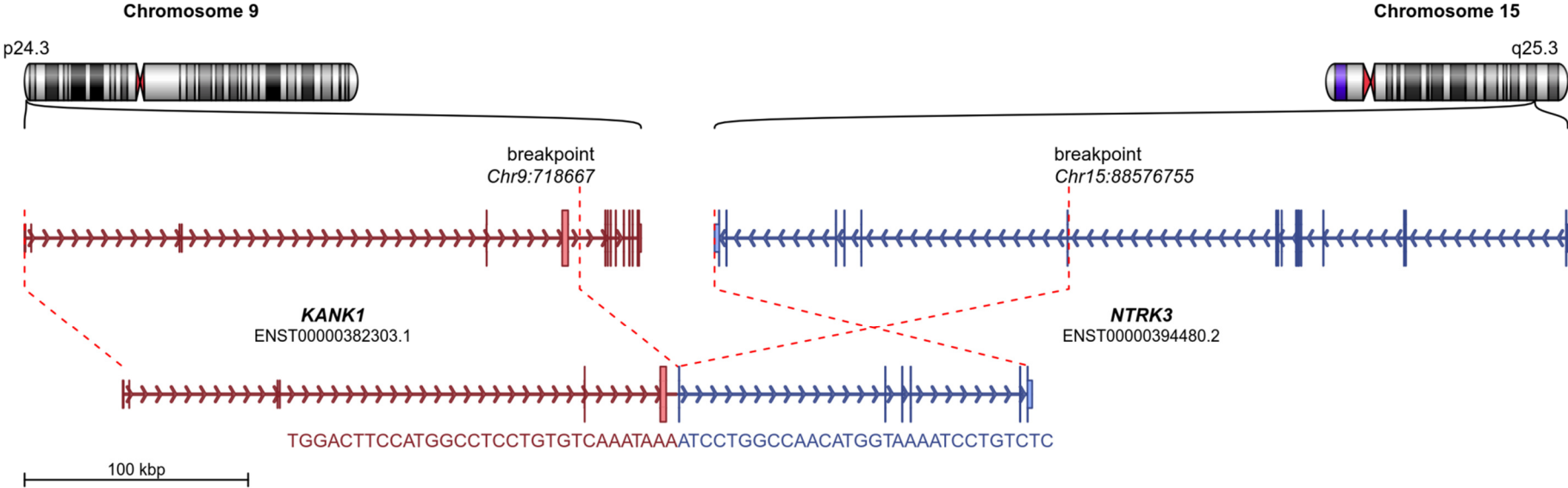
Patient A (EGAF00002486940)



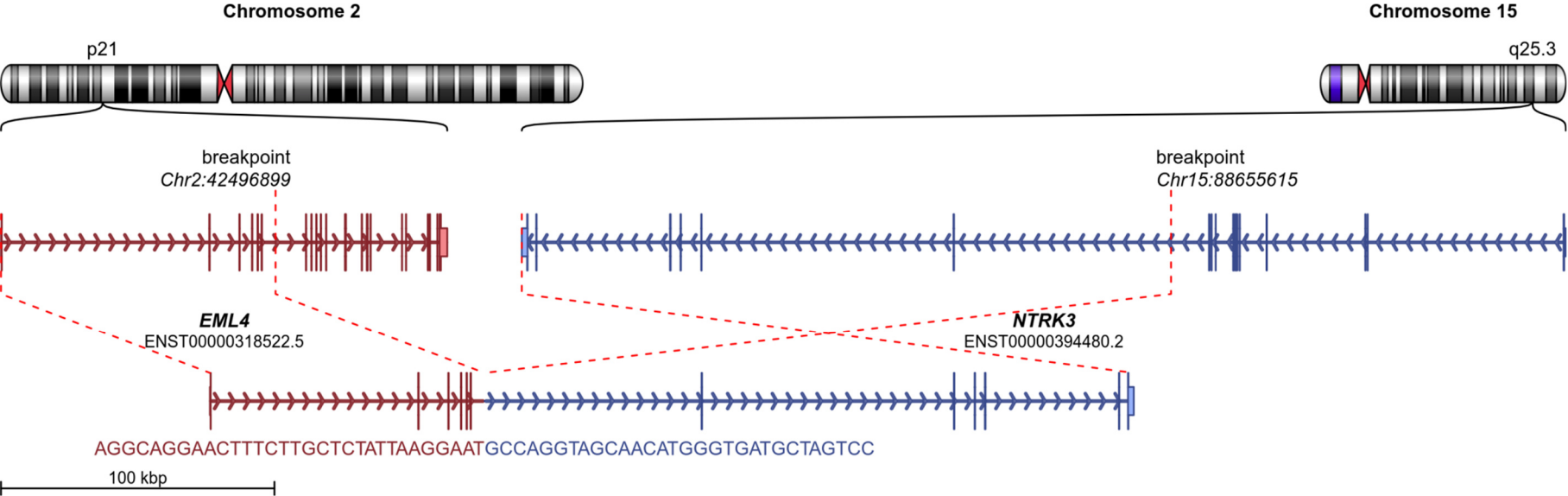
PCSI_0108 (EGAF00001556794)



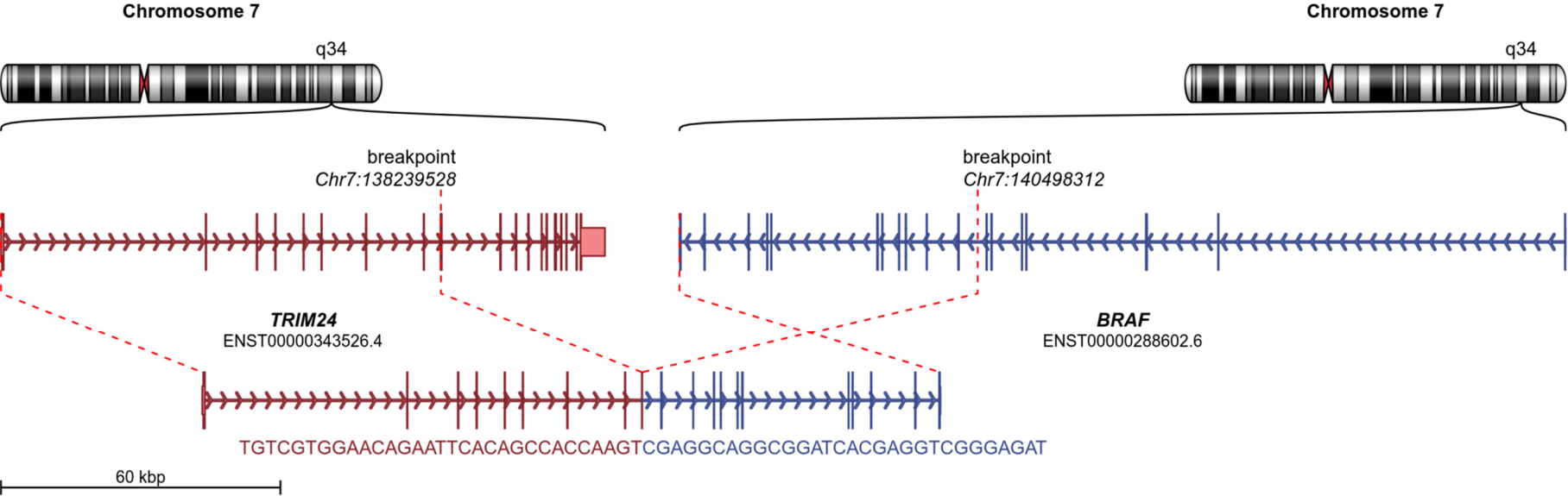
PCSI_0279 (EGAF00001556830)



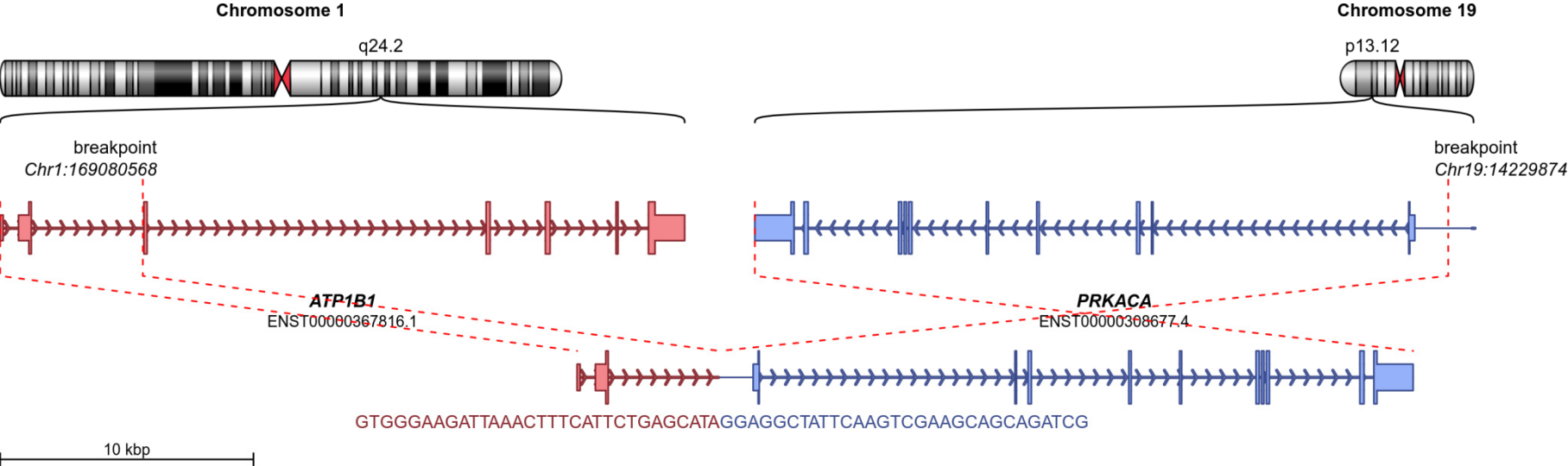
PCSI_0305 (EGAF00001556446)



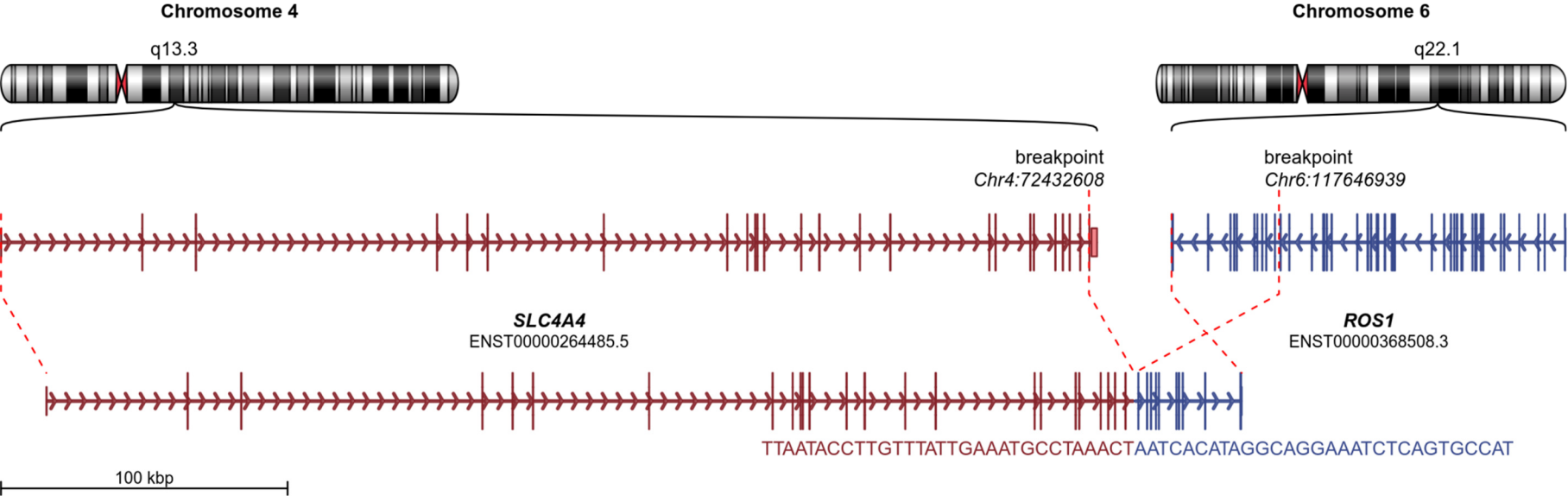
PCSI_0326 (EGAF00001556521)



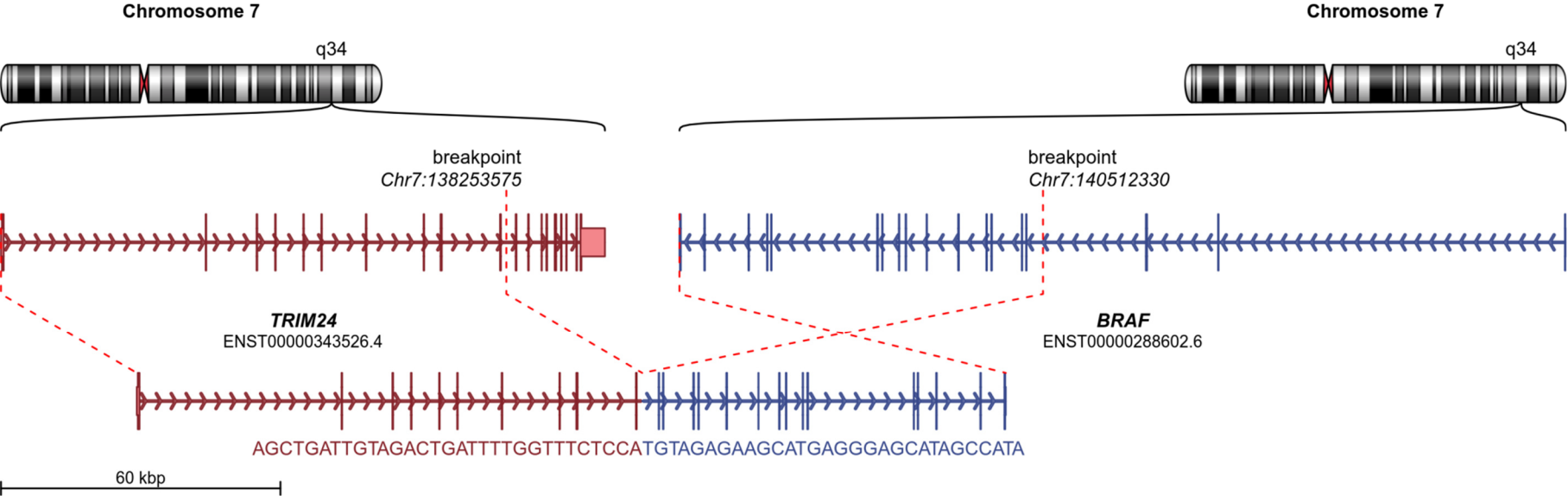
PCSI_0330 (EGAF00001802818)



PCSI_0347 (EGAF00001556815)



PCSI_0458 (EGAF00001556509)



PCSI_0572 (EGAF00001556614)

Chromosome 14

q32.32

Chromosome 2

p23.2

breakpoint
Chr14:103418480

breakpoint
Chr2:29446630



CDC42BPB
ENST00000381246.2

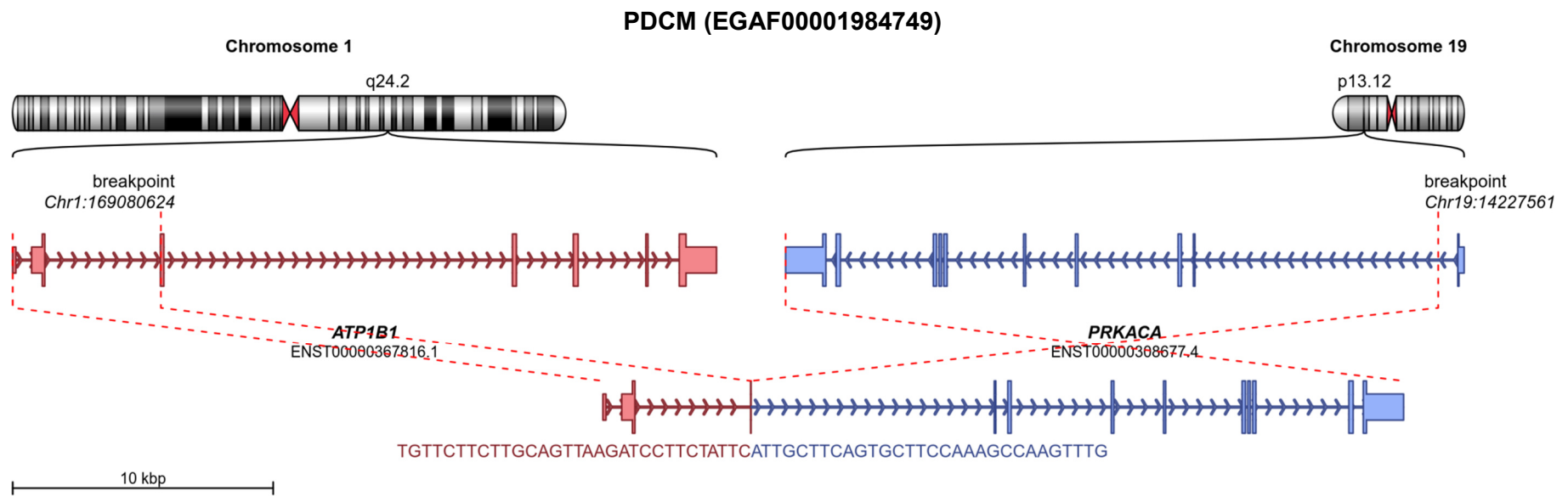


ALK
ENST00000389048.3



GATCTCACTTTGTTTCCCAGACTGGTCTCAATACAAATAATTATTTCCATATATCTGATT

200 kbp



Supplemental Figure S7: Structural variants correlating with the putative driver fusions in Supplemental Figure S6.
Structural variants were identified by SOPHIA in matched whole-genome sequencing samples.